

Sub
756

Notch(C)	IDEÇ-SNP	ÇONGGTÇ---	D-VGSY-Ç-ÇPPGFT	GK---ÇE-N
10244(C)	-NEÇTM---	ÇQH---	VNT-ÇSY-ÇKÇ-SÇ--	Ç--L-Ç D
80	VNEÇGMKPRP	ÇQHR Ç	ÇRÇFPÇYT	GKT ÇSQ D
95	VNSRTCAMIN	ÇQYS Ç	VNTHÇSYKÇFÇLS	GUMMP D
133	IDECASGKVI	ÇPYNRRC	EDTEEGPQÇLÇPSS	ÇLRLAPN
175	INEÇTMDSHT	ÇSHHANÇ	VNTFGSYVÇKÇHÇEE	LQYISGR
220			FNTQÇSF ÇKÇVÇYK	ÇNGRLÇS
ÇD27(C)	V-EÇ-SG-Q--Ç-SS--Ç	-NTVGSY-ÇRÇRPÇW-P-PÇ-PN---		D
EGF(C)	NSDSEÇPLSHDGYÇLJHDGVÇMYTEALDKYACNÇVVGYYI---	ÇER--ÇQYRDLKWWELR		

Figure 1

GGCTGGAGAA GAAACAGCAA GGGAGTCTGT GAAGCTACAT GCGAACTGG
ATGTAAGTTT GGTGAGTGCG TGGGACCAAA CAAATGCAGA TGCTTTCCAG
GATACACCGG GAAAACCTGC AGTCAAGATG TGAATGAGTG TGGAAAGAAA
CCCCGGCCAT GCCAACACAG ATGTGTGAAT ACACACGGAA GCTACAAGTG
CTTTTGCTC AGTGGCCACA TGCTCATGCC AGATGCTACG TGTGTGAAGT
CNAGGACATG TGCCATGATA AACTGTCAGT ATAGCTGTGA AGACACAGAA
(SEQ ID NO 1)

GGCTGGAGAA GAAACAGCAA GGGAGTCTGT GAAGCTACAT GCGAACCTGG
ATGTAAGTTT GGTGAGTGCG TGGGACCAAA CAAATGCAGA TGCTTTCCAG
GATACACCGG GAAAACCTGC AGTCAAGATG TGAATGAGTG TGGAAAGAAA
CCCCGGCCAT GCCAACACAG ATGTGTGAAT ACACACGGAA GCTACAAGTG
CTTTTGCTC AGTGGCCACA TGCTCATGCC AGATGCTACG TGTGTGAAGT
CNAGGACATG TGCCATGATA AACTGTCAGT ATAGCTGTGA AGACACAGAA
GAAGGGCCAC AGTGCCTGTG TCCATCCTCA GGAATCCGCC TGGCCCCAAA
TGGAAGAGAC TGTCTAGATA TTGATGAATG TGCCTCTGGT AAAGTCATCT
GTCCCTACAA TCGAAGATGT GTGAACACAT TTGGAAGCTA CTACTGCAAA
TGTCACATTG GTTTCGAACT GCAATATATC AGTGGACGAT ATGACTGTAT
AGATATAAAT GAATGTACTA TGGATAGCCA TACGTGCAGC CACCATGCCA
ATTGCTTCAA TACCCAAGGG TCCTTCAAGT GTAAATGCAA GCAGGGATAT
AAAGGCAATG GACTTCGGTG TTCTGCTATC CCTGAAAATT CTGTGAAGGA
AGTCCTCAGA GCACCTGGTA CCATCAAAGA CAGAATCAAG AAGTTGCTTG
CTCACAAAAA CAGCATGAAA AAGAAGGCAA AAATTAAAAA TGTTACCCCA
GAACCCACCA GGAATCCTAC CCCTAAGGTG AACTTGCAGC CCTTCAACTA
TGAAGAGATA GTTTCAGAG GCGGGAATC TCATGGAGGT AAAAAAGGGA
ATGAAGAGAA AATGAAAGAG GGGCTTGAGG ATGAGAAAAG AGAAGAGAAA
GCCCTGAAGA ATGACATAGA GGAGCGAAGC CTGCGAGGAG ATGTGTTTTT
CCCTAAGGTG AATGAAGCAG GTGAATTCCG CCTGATTCTG GTCCAAAGGA
AAGCGTAAC TTCCAACTG GAACATAAAG ATTTAAATAT CTCGGTGGAC
TGCAGCTTCA ATCATGGGAT CTGTGACTGG AAACAGGATA GAGAAGATGA
TTTTGACTGG AATCCTGCTG ATCGAGATAA TGCTATTGGC TTCTATATGG
CAGTTCCGGC CTTGGCAGGT CACATGAAAG ACATTGGCCG ATTGAAACTT
CTCCTACCTG ACCTGCAACC CCAAAGCAAC TTCTGTTTGC TCTTTGATTA
CCGGCTGGCC GGAGACAAAG TCGGGAAACT TCGAGTGTTT GTGAAAACA
GTAACAATGC CCTGGCATGG GAGAAGACCA CGAGTGAGGA TGAAAAGTGG
AAGACAGGGA AAATTCAATT GTATCAAGGA ACTGATGCTA CCAAAAGCAT
CATTTTGAAG GCAGAACGTG GCAAGGGCAA AACCAGCGAA ATCGCAGTGG
ATGGCGTCTT GCTTGTTC GGTATATGTC CAGATAGCCT TTTATCTGTG
GANNNCTGAA TGGTACTATC TTTATATTTG ACTTTGTATG TCAGTTCCCT
GGTTTTTTTG ATATTGCATC ATAGGACCTC TGGCATTTTA AAATTACTAG
CTGAAAAATTG
(SEQ ID NO 2)

Figure 2

GWRRNSKGVCEATCEPGCKFGECVGPNNKRCFPGYTGKTCSQDVNECGMKPRPCQHR
CVNTHGSYKCFCLSGHMLMPDATCVNSRTCAMINCQYSCEDTE
(SEQ ID NO 3)

GWRRNSKGVCEATCEPGCKFGECVGPNNKRCFPGYTGKTCSQDVNECGMKPRPCQHR
CVNTHGSYKCFCLSGHMLMPDATCVNSRTCAMINCQYSCEDTEEGPQCLCPSSGLRLAP
NGRDCLDIDECASGKVICPYNRRVCNTEFGSYCKCHIGFELQYISGRYDCIDINECTMDS
HTCSHHANCFNTQGSFKCKCKQGYKGNGLRCSAIPENSVKEVLRAPGTIKDRIKKLLAH
KNSMKKKAKIKNVTPEPTRTPKVNLPFNYYEIVSRGGNSHGGKKGNEEKMKEGLE
DEKREEKALKNDIEERSLRGDVFFPKVNEAGEFGLILVQRKALTSKLEHKDLNISVDCSF
NHGICDWKQDREDDFDWNPADRDNAIGFYMAVPALAGHMKDIGRLKLLLPDLQPQSN
FCLLFDYRLAGDKVGKLRVFKNSNNALAWKTTSEDEKWKTGKIQLYQGTDATKSHF
EAERGKGKTGEIAVDGVLLVSGLCPSLLSVDDXMVLSLYLTLYVSSLVFLILHRTSGI
LKLLAEKL
(SEQ ID NO 4)

Figure 3

09687860-10300

00000000000000000000000000000000

(SEQ ID NO: 5)

Figure 5

(SEQ ID NO: 6)

(SEQ ID NO: 6)

EGFL6 (221-260 aa)
3D Model

EGF
NMR Structure

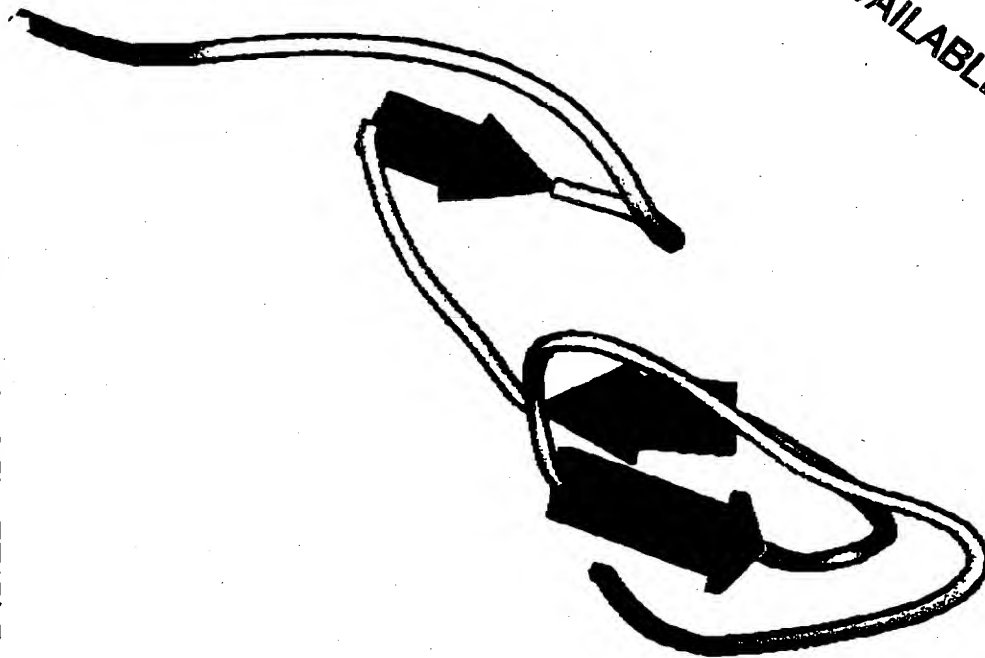
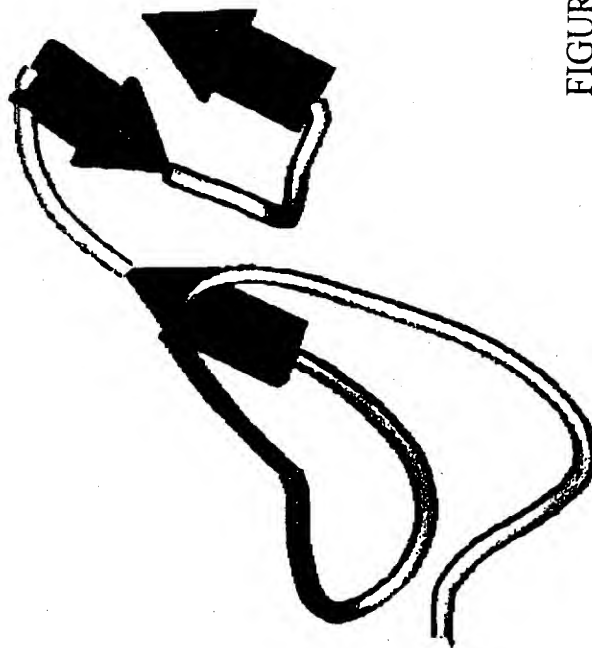


FIGURE 6

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